

PEPTIDESTRUCTURE of: human\_prot.pep ck: 2321, 1 to 458

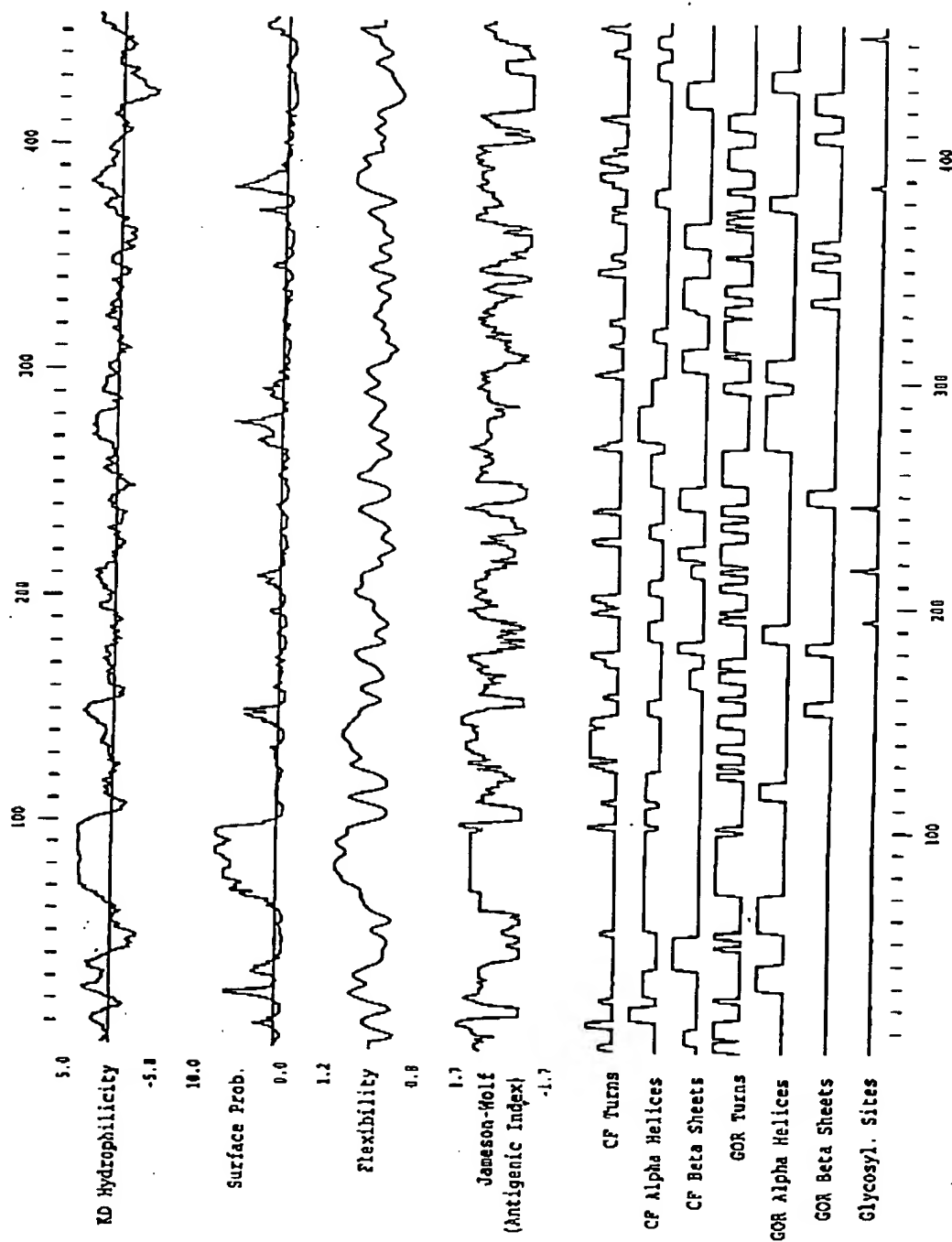
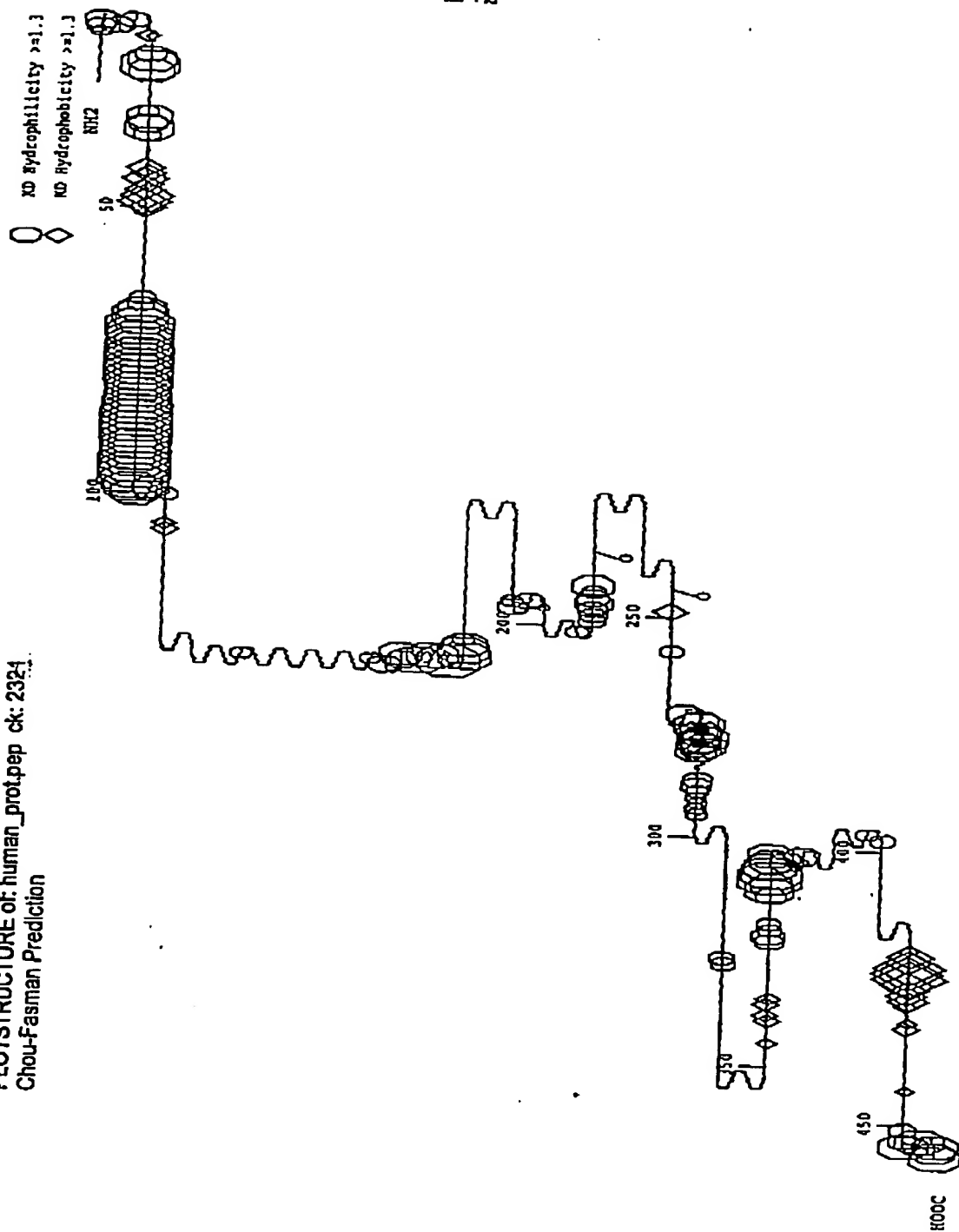


FIGURE 1

662TQT\*HBE9TH60

PLOTSTRUCTURE of: human\_prot.pep ck: 2324  
Chou-Fasman Prediction



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FIGURE 2

662TQT"48E9T460

PLOTSTRUCTURE of: human\_prot.pep ck: 2321.  
Garner-Osguthorpe-Robson Prediction

○ Hydrophilicity >1.3  
◇ Hydrophobicity >1.3

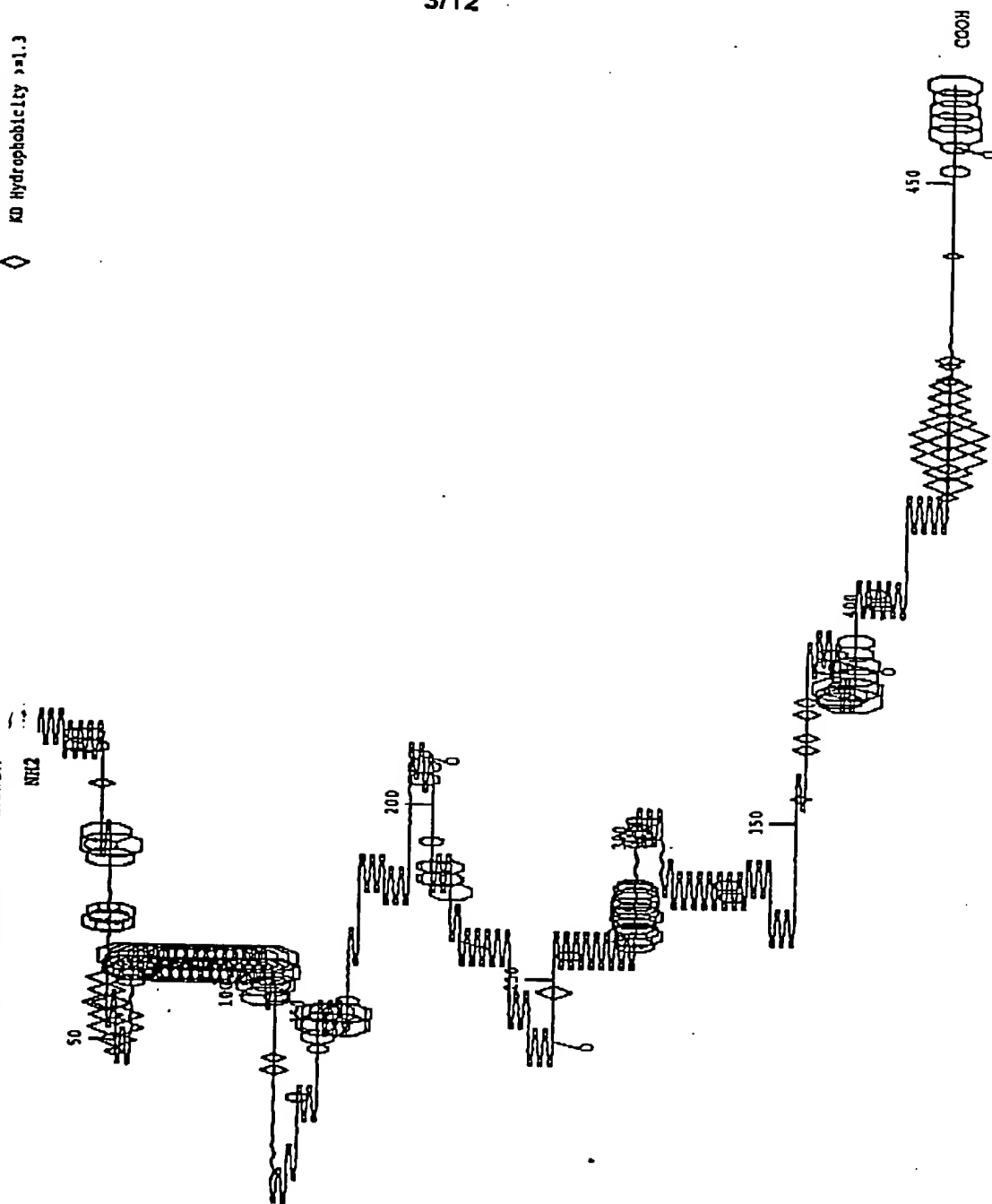


FIGURE 3

PEPTIDESTRUCTURE of: g713estmouse.0.12\_tr13.pap ck: 1880, 1 to 467

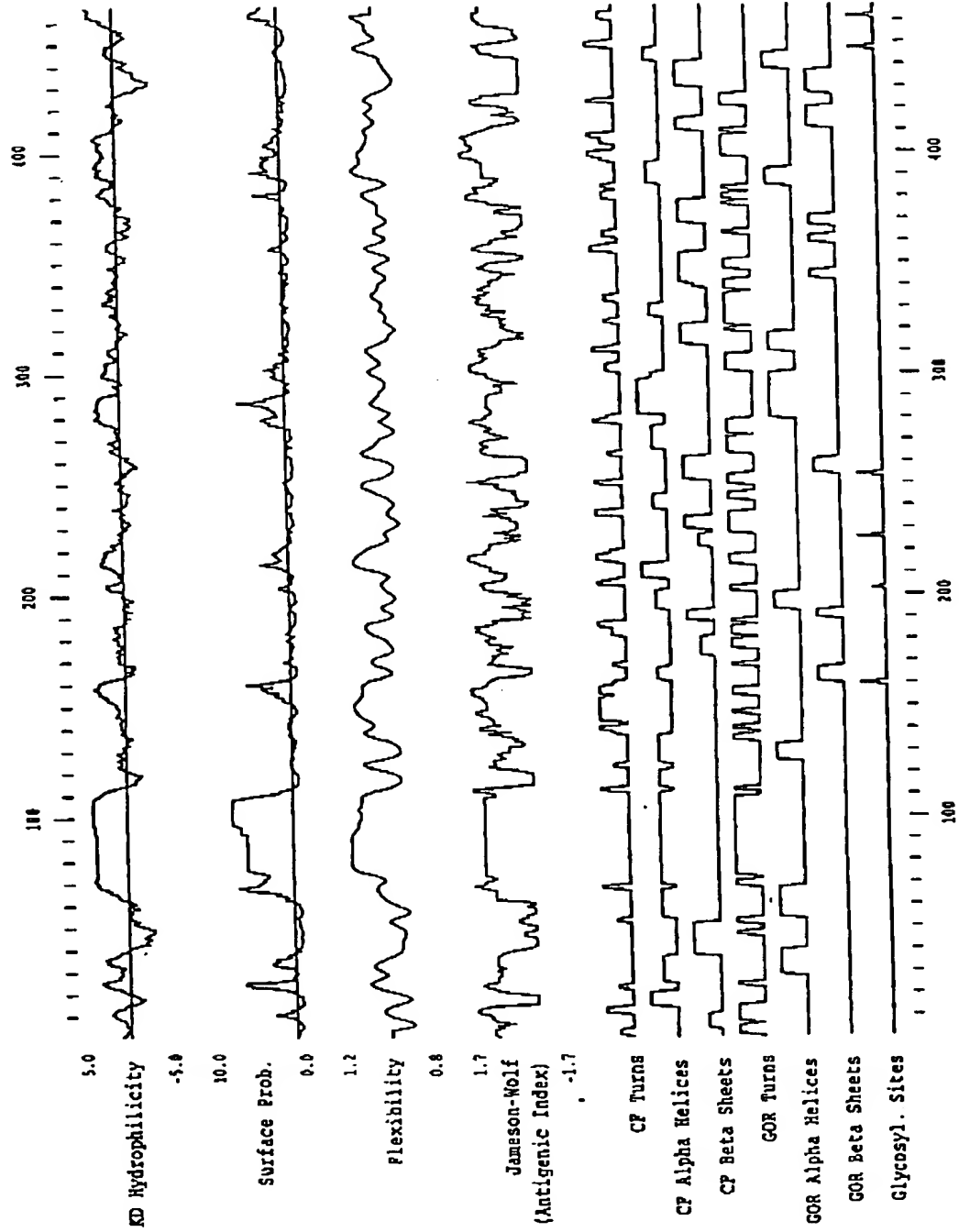
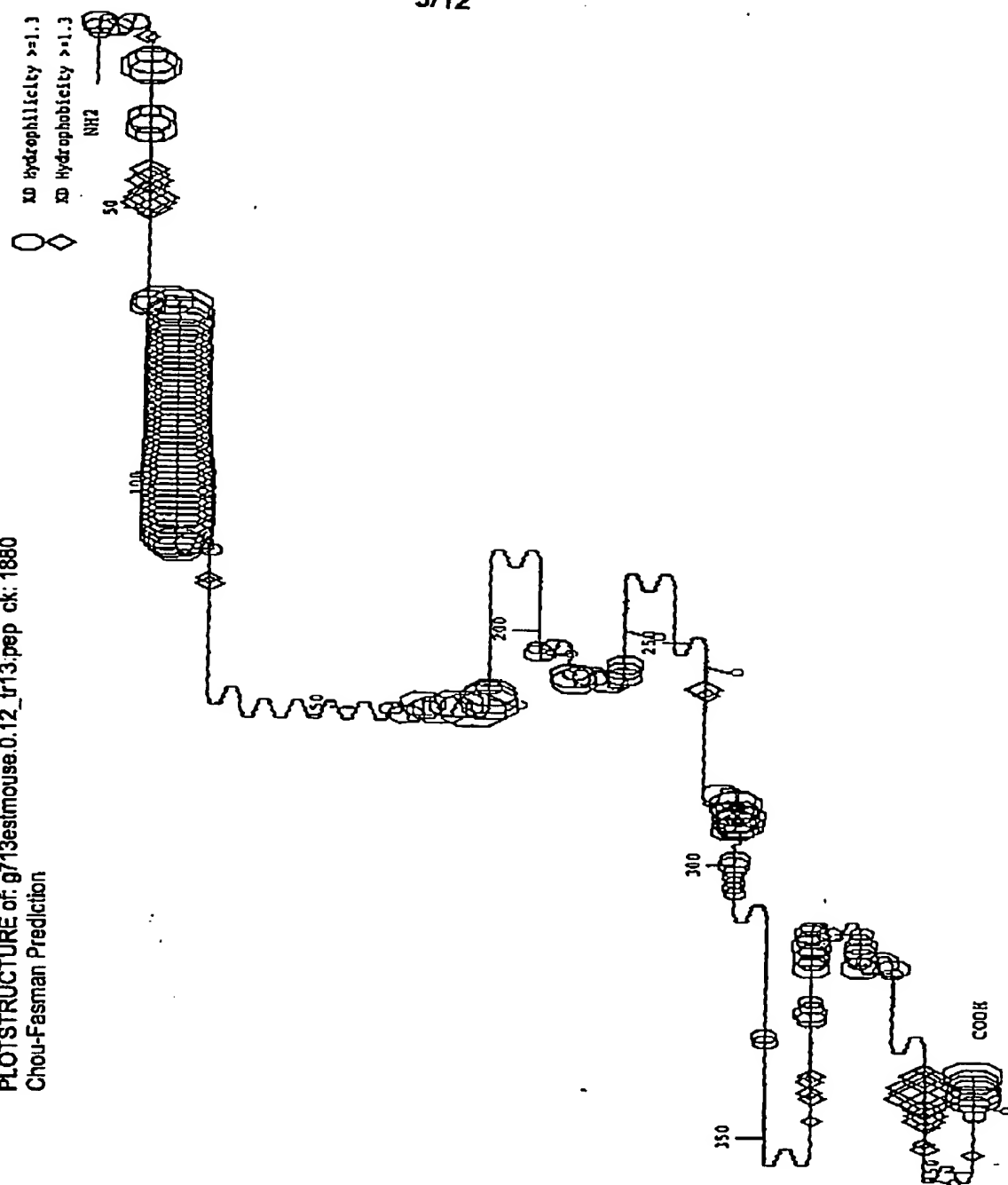


FIGURE 4

662T0T"4BESF460

PLOTSTRUCTURE of: g713estmouse.0.12\_tr13.pep ck: 1880  
Chou-Fasman Prediction



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FIGURE 5

662TQT"48E9T460

PLOTSTRUCTURE of: g713estmouse.0.12\_tr13.pép ck: 1880  
Garnier-Osguthorpe-Robson Prediction

XD Hydrophilicity >=1.3  
XD Hydrophobicity >=1.3

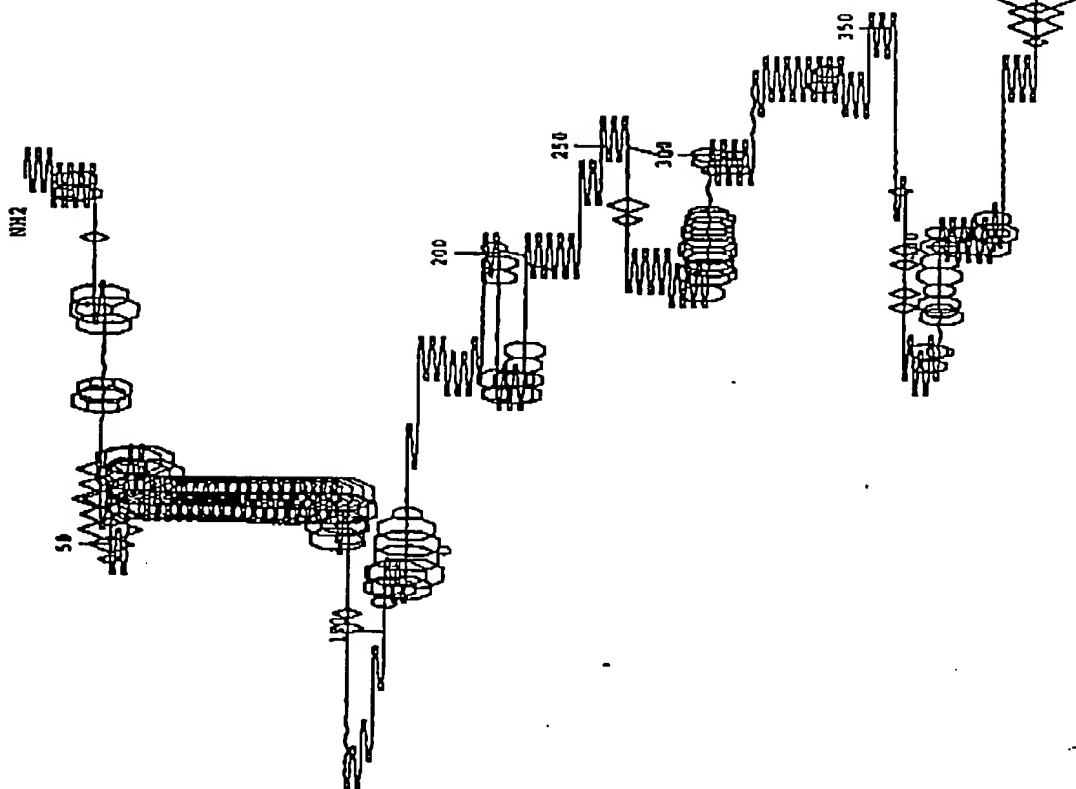


FIGURE 6

09416384.101299

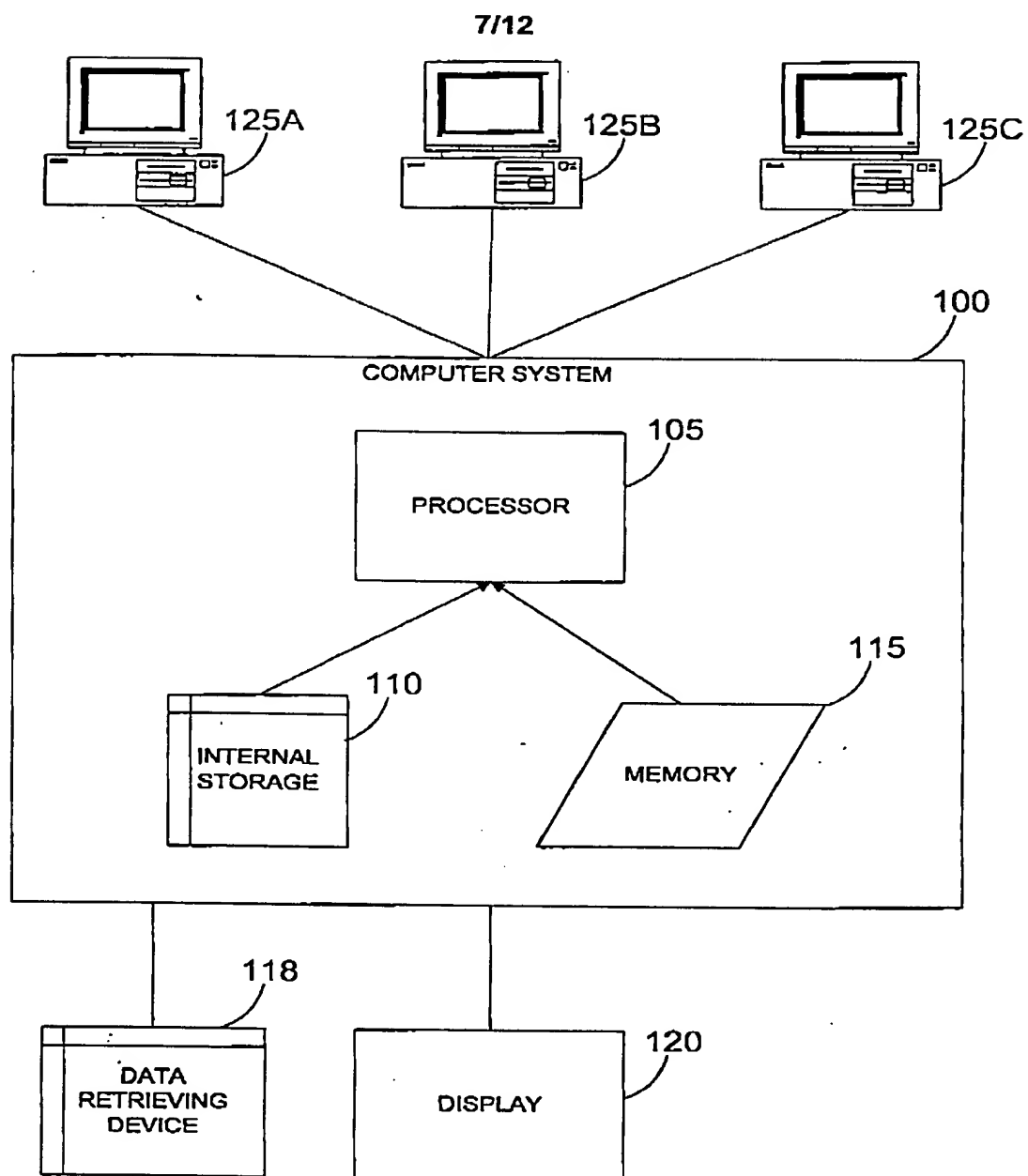


FIGURE 7

09416384-101299

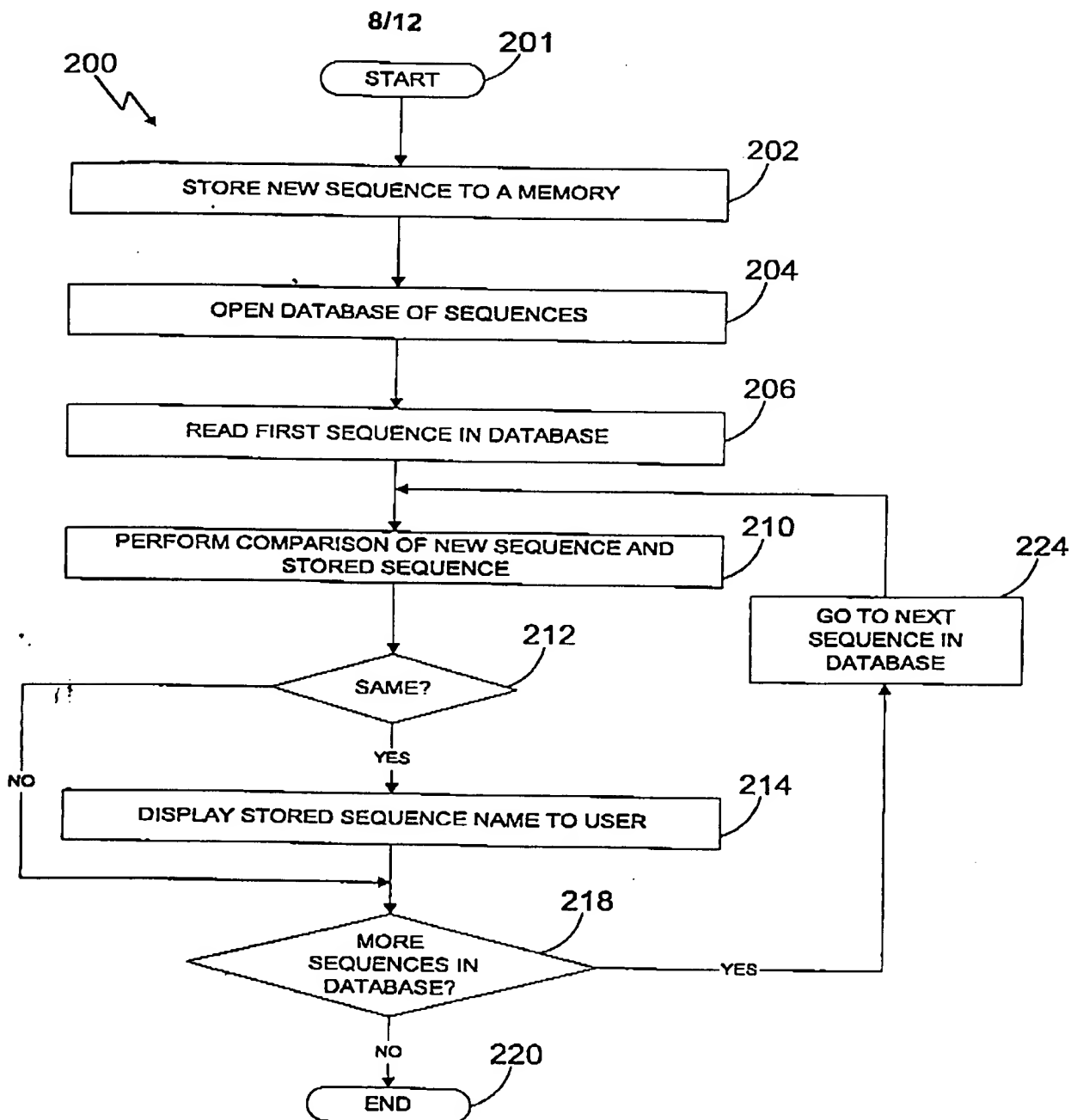


FIGURE 8



00416384-101299

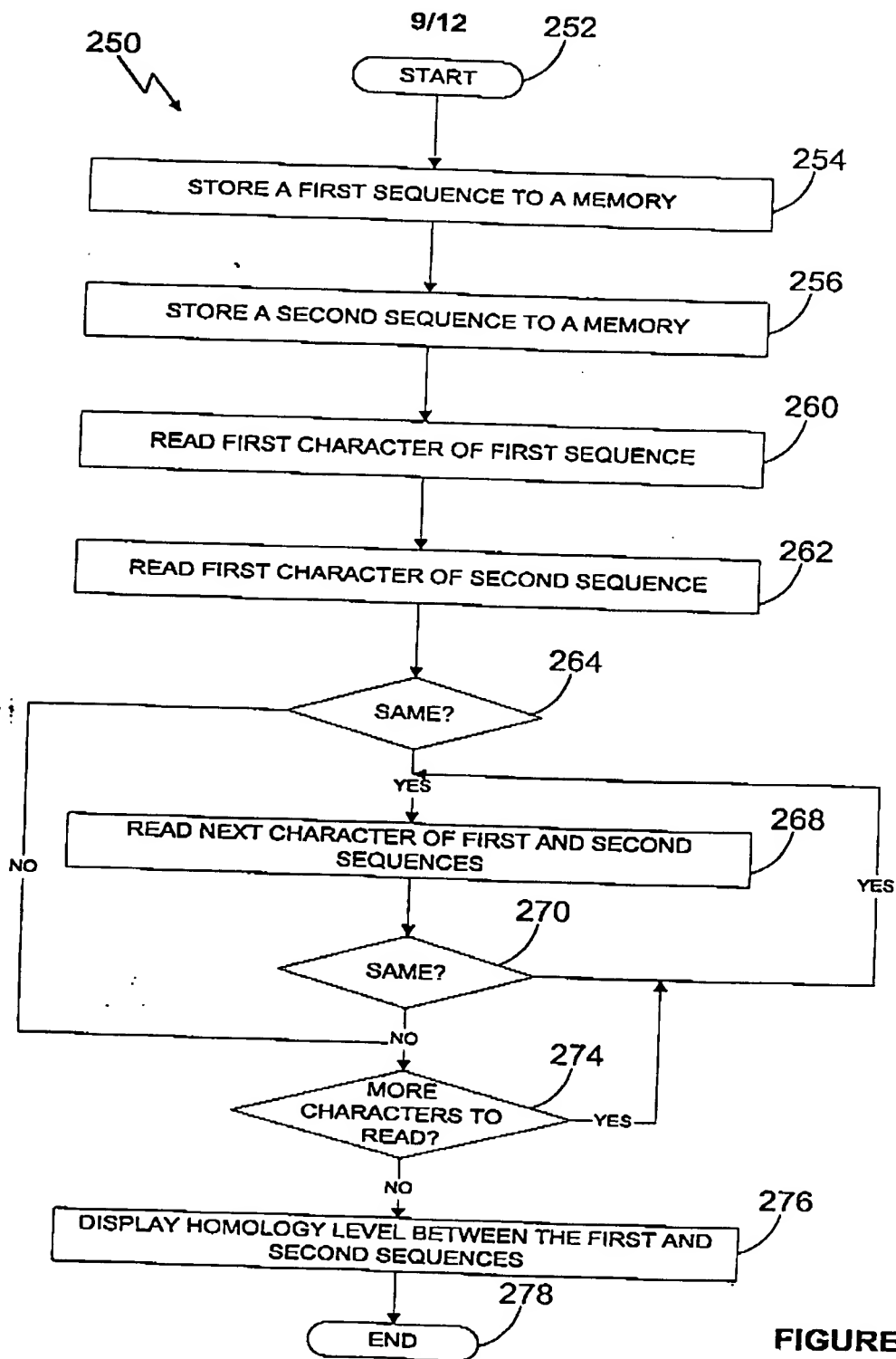


FIGURE 9

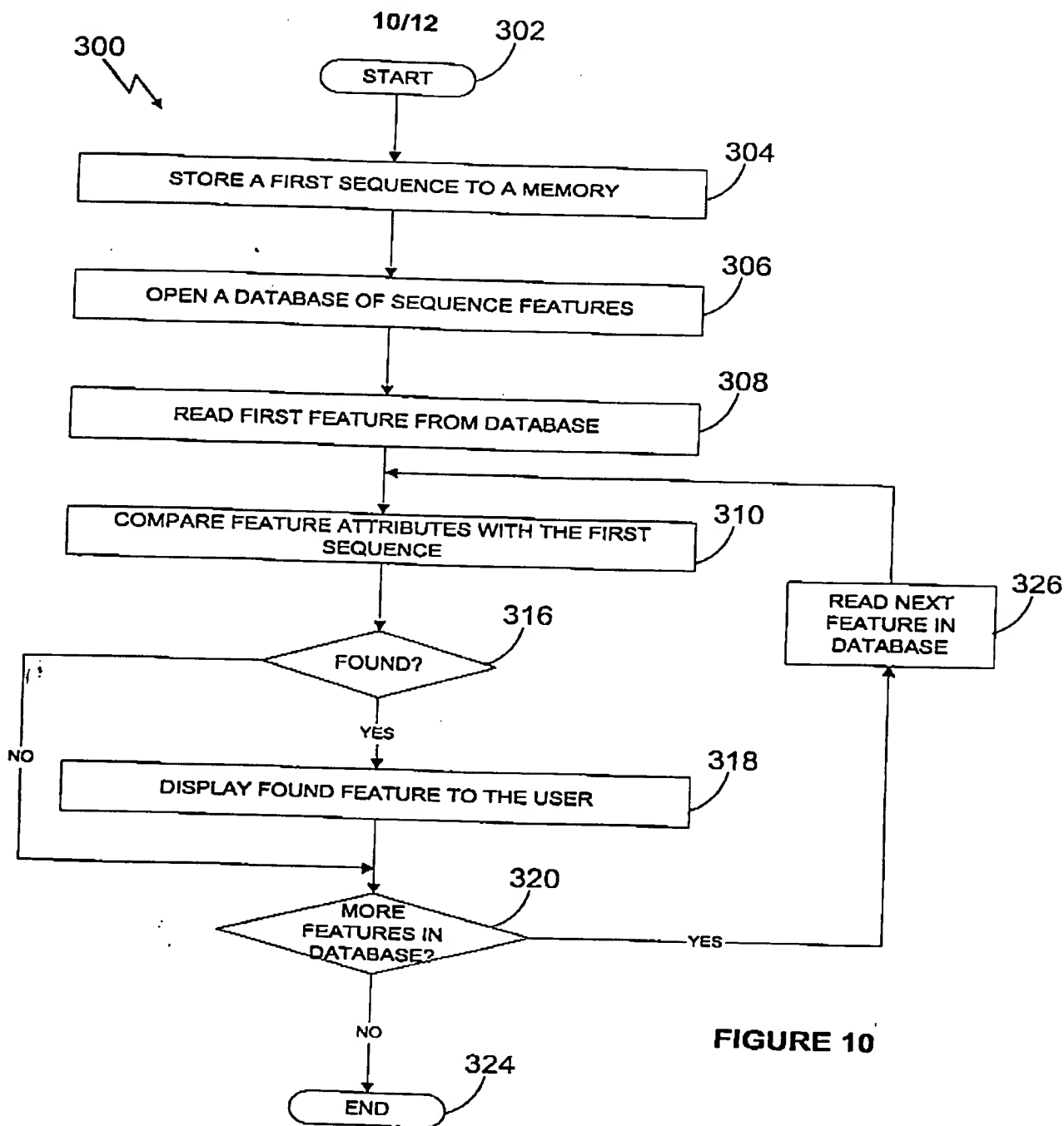


FIGURE 10

FIGURE 11

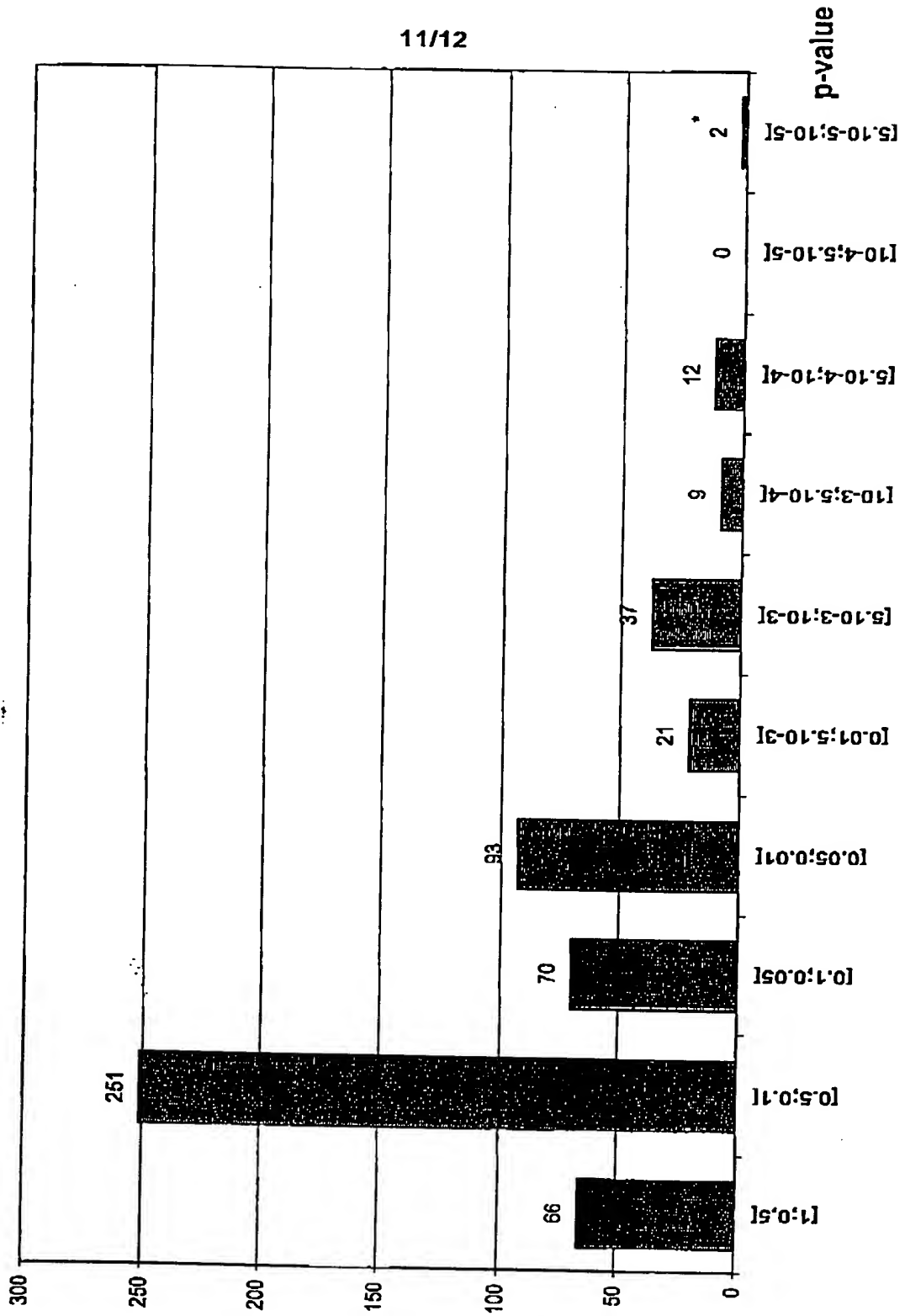


FIGURE 12

